UY//U1245

WO 99/61661

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PCT/FR99/01247

JC01 Rec'd PCT/PTO 2 7 NOV 200

SEQUENCE LISTING

- (1) GENERAL INFORMATION:
 - (i) APPLICANT:
 - (A) NAME: BIOMERIEUX
 - (B) STREET: CHEMIN DE L'ORME
 - (C) CITY: MARCY L'ETOILE
 - (E) COUNTRY: FRANCE
 - (F) POSTAL CODE: 69280
 - (ii) TITLE OF THE INVENTION: Method for amplifying at least one specific nucleotide sequence, and primers used
- (iii) NUMBER OF SEQUENCES: 19
- (iv) CORRESPONDENCE ADDRESS:
 - (A) NAME: Cabinet GERMAIN & MAUREAU
 - (B) STREET: 12 rue Boileau
 - (C) CITY: LYON
 - (D) COUNTRY: FRANCE
 - (E) POSTAL CODE: 69006
 - (F) TELEPHONE: 04 72 69 84 30
 - (G) TELEPAX: 04 72 69 84 31
 - (v) COMPUTER READABLE FORM
 - (A) MEDIUM TYPE: DS, HD 3.5 inch disk
 - (B) COMPUTER: EASON (IBM compatible)
 - (C) OPERATING SYSTEM: DOS
 - (D) SOFTWARE: MICROSOFT WORD FOR WINDOWS
- (2) INFORMATION FOR SEQ ID NO: 1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 23 base pairs
 - (B) TYPE: nucleotide
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear



(ii)	MOLECULE TYPE: cDNA	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 1:	
ATCCTTC	CGTG TCCCCACAGC ACG	23
2) INFO	ORMATION FOR SEQ ID NO: 2:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 19 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 2:	
TCGCCGC	TGC ACTGTGAAG	19
2) INFO	DRMATION FOR SEQ ID NO: 3:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 24 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(ix)	CHARACTERISTIC:	
	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 24 (3' end)	
	(D) OTHER INFORMATION: modification by the [lacuna] group	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 3:	
CCCCCA	GCA CGTTTCTTGG AGCT 24	
(2) INFO	DRMATION FOR SEQ ID NO: 4:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 24 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	

(ix)	CHARACTERISTIC:	
	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 1 (5' end)	
	(D) OTHER INFORMATION: modification by the acridine group	qı
(ix)	CHARACTERISTIC:	
	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 24 (3' end)	
	(D) OTHER INFORMATION: modification by H	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 4:	
CCCCCA	GCA CGTTTCTTGG AGCT	24
(2) INFO	ORMATION FOR SEQ ID NO: 5:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 26 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(ix)	CHARACTERISTIC:	
	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 24	
	(D) OTHER INFORMATION: G means inosine (i)	
(ix)	CHARACTERISTIC:	
	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 26 (3' end)	
	(D) OTHER INFORMATION: modification by the C_6-NH_2 group	
(xi)	7	26
CCCACAG	CAC GTTTCTTGGA GCAGGC	26
(2) INF	ORMATION FOR SEQ ID NO: 6:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 21 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	

(ii)	MOLECULE TYPE: cDNA	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 6:	
CCCAGCA	CGT TTCTTGGAGC T	21
(2) INF	ORMATION FOR SEQ ID NO: 7:	
(i)		
(-,	(A) LENGTH: 24 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(;;)	MOLECULE TYPE: cDNA	
(xi)		
, ,	GCA CGTTTCTTGG AGCT	24
(2) INF	ORMATION FOR SEQ ID NO: 8:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 24 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ix)	CHARACTERISTIC:	
	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 23	
	(D) OTHER INFORMATION: G means inosine (i)	
(ii)	MOLECULE TYPE: cDNA	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 8:	
CCCCCF	AGCA CGTTTCTTGG AGGT	24
(2) INF	ORMATION FOR SEQ ID NO: 9:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 24 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)		
, ,	CHARACTERISTIC.	

	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 22	
	(D) OTHER INFORMATION: G means inosine (i)	
(ix)	CHARACTERISTIC:	
	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 23	
	(D) OTHER INFORMATION: G means inosine (i)	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 9:	
CATTTCC	TCA ATGGGACGGA GGGA	24
(2) INF	ORMATION FOR SEQ ID NO: 10:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 26 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(ix)	CHARACTERISTIC:	
	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 24	
	(D) OTHER INFORMATION: G means inosine (i)	
(:	xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:	
CCCCCAG	CAC GTTTCTTGGA GCAGGC	26
(2) INF	ORMATION FOR SEQ ID NO: 11:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 26 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(ix)	CHARACTERISTIC:	
	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 24	
	(D) OTHER INFORMATION: G means inosine (i)	
(201)	SECTIONS DESCRIPTION. SEC ID NO. 11.	

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CCCACAGCAC GTTTCTTGGA GCAGGC

(2) INF	ORMATION FOR SEQ ID NO: 12:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 19 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 12:	
CACGTTT	CTT GCAGCAGGA	19
(2) INF	ORMATION FOR SEQ ID NO: 13:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 22 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 13:	
CAGCACO	TTT CTTGCAGCAG GA	22
(2) INF	ORMATION FOR SEQ ID NO: 14:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 19 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(ix)	CHARACTERISTIC:	
	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 3	
	(D) OTHER INFORMATION: G means inosine (i)	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 14:	
CAGGTTT	CTT GCAGCAGGA	19

(2) INF	ORMATION FOR SEQ ID NO: 15:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 22 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(ix)	CHARACTERISTIC:	
	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 6	
	(D) OTHER INFORMATION: G means inosine (i)	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 15:	
CAGCAGG	TTT CTTGCAGCAG GA	22
(2) INF	ORMATION FOR SEQ ID NO: 16:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 26 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(ix)	CHARACTERISTIC:	
	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 10	
	(D) OTHER INFORMATION: G means inosine (i)	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 16:	
CCCCCAG	CAG GTTTCTTGCA GCAGGA	26
(2) INF	ORMATION FOR SEQ ID NO: 17:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 26 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(ix)	CHARACTERISTIC:	

	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 10	
	(D) OTHER INFORMATION: G means inosine (i)	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 17:	
CCCACAC	GCAG GTTTCTTGCA GCAGGA	26
(2) INF	ORMATION FOR SEQ ID NO: 18:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 26 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(ix)	CHARACTERISTIC:	
	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 10	
	(D) OTHER INFORMATION: G means inosine (i)	
(ix)	CHARACTERISTIC:	
	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 25	
	(D) OTHER INFORMATION: G means inosine (i)	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 18:	
CCCACAC	CAG GTTTCTTGCA GCAGGA	26
(2) INF	ORMATION FOR SEQ ID NO: 19:	
(i)	SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 26 base pairs	
	(B) TYPE: nucleotide	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(ix)	CHARACTERISTIC:	
	(A) NAME/KEY: particular characteristic	
	(B) LOCATION: 10	
	(D) OTHER INFORMATION: G means inosine (i)	
(ix)	CHARACTERISTIC:	

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(A) NAME/KEY: particular characteristic

(B) LOCATION: 25

(D) OTHER INFORMATION: G means inosine (i)

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:

CCCCCAGCAG GTTTCTTGCA GCAGGA

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